

SEQUENCE LISTING

<110> Boehringer Ingelheim Vetmedica GmbH

<120> Recombinant attenuation of Porcine Reproductive and Respiratory Syndrome (PRRSV)

<130> 1-1120,25,26-1-C1

<140> to be assigned
<141> June 20, 2003

<150> 09/772,316

<151> January 26, 2001

<150> DE 10003371.7

<151> 2000-01-26

<150> DE 10003373.3

<151> 2000-01-26

<150> DE 10003372.5

<151> 2000-01-26

<160> 6

<170> PatentIn Ver. 2.1

<210> 1

<211> 2502

<212> PRT

<213> Porcine reproductive and respiratory syndrome virus

<400> 1

Met Ser Gly Ile Leu Asp Arg Cys Thr Cys Thr Pro Asn Ala Arg Val
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Phe Met Ala Glu Gly Gln Val Tyr Cys Thr Arg Cys Leu Ser Ala Arg
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Ser Leu Leu Pro Leu Asn Leu Gln Val Ser Glu Leu Gly Val Leu Gly
35 40 45

Leu Phe Tyr Arg Pro Glu Glu Pro Leu Arg Trp Thr Leu Pro Arg Ala
50 55 60

Phe Pro Thr Val Glu Cys Ser Pro Ala Gly Ala Cys Trp Leu Ser Ala
65 70 75 80

Ile Phe Pro Ile Ala Arg Met Thr Ser Gly Asn Leu Asn Phe Gln Gln
85 90 95

Arg Met Val Arg Val Ala Ala Glu Leu Tyr Arg Ala Gly Gln Leu Thr
100 105 110

Pro Ala Val Leu Lys Ala Leu Gln Val Tyr Glu Arg Gly Cys Arg Trp
115 120 125

Tyr Pro Ile Val Gly Pro Val Pro Gly Val Ala Val Phe Ala Asn Ser
130 135 140

Leu His Val Ser Asp Lys Pro Phe Pro Gly Ala Thr His Val Leu Thr
145 150 155 160

Asn Leu Pro Leu Pro Gln Arg Pro Lys Pro Glu Asp Phe Cys Pro Phe
165 170 175

Glu Cys Ala Met Ala Thr Val Tyr Asp Ile Gly His Asp Ala Val Met
180 185 190

Tyr Val Ala Glu Arg Lys Val Ser Trp Ala Pro Arg Gly Gly Asp Glu
195 200 205

Val Lys Phe Glu Ala Val Pro Gly Glu Leu Lys Leu Ile Ala Asn Arg
210 215 220

Leu Arg Thr Ser Phe Pro Pro His His Thr Val Asp Met Ser Lys Phe
225 230 235 240

Ala Phe Thr Ala Pro Gly Cys Gly Val Ser Met Arg Val Glu Arg Gln
245 250 255

His Gly Cys Leu Pro Ala Asp Thr Val Pro Glu Gly Asn Cys Trp Trp
260 265 270

Ser Leu Phe Asp Leu Leu Pro Leu Glu Val Gln Asn Lys Glu Ile Arg
275 280 285

His Ala Asn Gln Phe Gly Tyr Gln Thr Lys His Gly Val Ser Gly Lys
290 295 300

Tyr Leu Gln Arg Arg Leu Gln Val Asn Gly Leu Arg Ala Val Thr Asp
305 310 315 320

Leu Asn Gly Pro Ile Val Val Gln Tyr Phe Ser Val Lys Glu Ser Trp
325 330 335

Ile Arg His Leu Lys Leu Ala Gly Glu Pro Ser Tyr Ser Gly Phe Glu
340 345 350

Asp Leu Leu Arg Ile Arg Val Glu Pro Asn Thr Ser Pro Leu Ala Asp
355 360 365

Lys Glu Glu Lys Ile Phe Arg Phe Gly Ser His Lys Trp Tyr Gly Ala
370 375 380

Gly Lys Arg Ala Arg Lys Ala Arg Ser Cys Ala Thr Ala Thr Val Ala
385 390 395 400

Gly Arg Ala Leu Ser Val Arg Glu Thr Arg Gln Ala Lys Glu His Glu
405 410 415

Val Ala Gly Ala Asn Lys Ala Glu His Leu Lys His Tyr Ser Pro Pro
420 425 430

Ala Glu Gly Asn Cys Gly Trp His Cys Ile Ser Ala Ile Ala Asn Arg
435 440 445

Met Val Asn Ser Lys Phe Glu Thr Thr Leu Pro Glu Arg Val Arg Pro
450 455 460

Pro Asp Asp Trp Ala Thr Asp Glu Asp Leu Val Asn Ala Ile Gln Ile
465 470 475 480

Leu Arg Leu Pro Ala Ala Leu Asp Arg Asn Gly Ala Cys Thr Ser Ala
485 490 495

Lys Tyr Val Leu Lys Leu Glu Gly Glu His Trp Thr Val Thr Val Thr
500 505 510

Pro Gly Met Ser Pro Ser Leu Leu Pro Leu Glu Cys Val Gln Gly Cys
515 520 525

Cys Gly His Lys Gly Gly Leu Gly Ser Pro Asp Ala Val Glu Val Ser
530 535 540

Gly Phe Asp Pro Ala Cys Leu Asp Arg Leu Ala Glu Val Met His Leu
545 550 555 560

Pro Ser Ser Ala Ile Pro Ala Ala Leu Ala Glu Met Ser Gly Asp Ser
565 570 575

Asp Arg Ser Ala Ser Pro Val Thr Thr Val Trp Thr Val Ser Gln Phe
580 585 590

Phe Ala Arg His Ser Gly Gly Asn His Pro Asp Gln Val Arg Leu Gly
595 600 605

Lys Ile Ile Ser Leu Cys Gln Val Ile Glu Asp Cys Cys Ser Gln
610 615 620

Asn Lys Thr Asn Arg Val Thr Pro Glu Glu Val Ala Ala Lys Ile Asp
625 630 635 640

Leu Tyr Leu Arg Gly Ala Thr Asn Leu Glu Glu Cys Leu Ala Arg Leu
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Glu Lys Ala Arg Pro Pro Arg Val Ile Asp Thr Ser Phe Asp Trp Asp
660 665 670

Val Val Leu Pro Gly Val Glu Ala Ala Thr Gln Thr Ile Lys Leu Pro
675 680 685

Gln Val Asn Gln Cys Arg Ala Leu Val Pro Val Val Thr Gln Lys Ser
690 695 700

Leu Asp Asn Asn Ser Val Pro Leu Thr Ala Phe Ser Leu Ala Asn Tyr
705 710 715 720

Tyr Tyr Arg Ala Gln Gly Asp Glu Val Arg His Arg Glu Arg Leu Thr
725 730 735

Ala Val Leu Ser Lys Leu Glu Lys Val Val Arg Glu Glu Tyr Gly Leu
740 745 750

Met Pro Thr Glu Pro Gly Pro Arg Pro Thr Leu Pro Arg Gly Leu Asp
755 760 765

Glu Leu Lys Asp Gln Met Glu Glu Asp Leu Leu Lys Leu Ala Asn Ala
770 775 780

Gln Thr Thr Ser Asp Met Met Ala Trp Ala Val Glu Gln Val Asp Leu
785 790 795 800

Lys Thr Trp Val Lys Asn Tyr Pro Arg Trp Thr Pro Pro Pro Pro Pro
805 810 815

Pro Lys Val Gln Pro Arg Lys Thr Lys Pro Val Lys Ser Leu Pro Glu
820 825 830

Arg Lys Pro Val Pro Ala Pro Arg Arg Lys Val Gly Ser Asp Cys Gly
835 840 845

Ser Pro Val Ser Leu Gly Gly Asp Val Pro Asn Ser Trp Glu Asp Leu
850 855 860

Ala Val Ser Ser Pro Phe Asp Leu Pro Thr Pro Pro Glu Pro Ala Thr
865 870 875 880

Pro Ser Ser Glu Leu Val Ile Val Ser Ser Pro Gln Cys Ile Phe Arg
885 890 895

Pro Ala Thr Pro Leu Ser Glu Pro Ala Pro Ile Pro Ala Pro Arg Gly
900 905 910

Thr Val Ser Arg Pro Val Thr Pro Leu Ser Glu Pro Ile Pro Val Pro
915 920 925

Ala Pro Arg Arg Lys Phe Gln Gln Val Lys Arg Leu Ser Ser Ala Ala
930 935 940

Ala Ile Pro Pro Tyr Gln Asp Glu Pro Leu Asp Leu Ser Ala Ser Ser
945 950 955 960

Gln Thr Glu Tyr Glu Ala Ser Pro Pro Ala Pro Pro Gln Ser Gly Gly
965 970 975

Val Leu Gly Val Glu Gly His Glu Ala Glu Glu Thr Leu Ser Glu Ile
980 985 990

Ser Asp Met Ser Gly Asn Ile Lys Pro Ala Ser Val Ser Ser Ser Ser
995 1000 1005

Ser Leu Ser Ser Val Arg Ile Thr Arg Pro Lys Tyr Ser Ala Gln Ala
1010 1015 1020

Ile Ile Asp Ser Gly Gly Pro Cys Ser Gly His Leu Gln Glu Val Lys
1025 1030 1035 1040

Glu Thr Cys Leu Ser Val Met Arg Glu Ala Cys Asp Ala Thr Lys Leu
1045 1050 1055

Asp Asp Pro Ala Thr Gln Glu Trp Leu Ser Arg Met Trp Asp Arg Val
1060 1065 1070

Asp Met Leu Thr Trp Arg Asn Thr Ser Val Tyr Gln Ala Ile Cys Thr
1075 1080 1085

Leu Asp Gly Arg Leu Lys Phe Leu Pro Lys Met Ile Leu Glu Thr Pro
1090 1095 1100

Pro Pro Tyr Pro Cys Glu Phe Val Met Met Pro His Thr Pro Ala Pro
1105 1110 1115 1120

Ser Val Gly Ala Glu Ser Asp Leu Thr Ile Gly Ser Val Ala Thr Glu
1125 1130 1135

Asp Val Pro Arg Ile Leu Glu Lys Ile Glu Asn Val Gly Glu Met Ala
1140 1145 1150

Asn Gln Gly Pro Leu Ala Phe Ser Glu Asp Lys Pro Val Asp Asp Gln
1155 1160 1165

Leu Val Asn Asp Pro Arg Ile Ser Ser Arg Arg Pro Asp Glu Ser Thr
1170 1175 1180

Ser Ala Pro Ser Ala Gly Thr Gly Gly Ala Gly Ser Phe Thr Asp Leu
1185 1190 1195 1200

Pro Pro Ser Asp Gly Ala Asp Ala Asp Gly Gly Pro Phe Arg Thr
1205 1210 1215

Val Lys Arg Lys Ala Glu Arg Leu Phe Asp Gln Leu Ser Arg Gln Val
1220 1225 1230

Phe Asp Leu Val Ser His Leu Pro Val Phe Phe Ser Arg Leu Phe Tyr
1235 1240 1245

Pro Gly Gly Gly Tyr Ser Pro Gly Asp Trp Gly Phe Ala Ala Phe Thr
1250 1255 1260

Leu Leu Cys Leu Phe Leu Cys Tyr Ser Tyr Pro Ala Phe Gly Ile Ala
1265 1270 1275 1280

Pro Leu Leu Gly Val Phe Ser Gly Ser Ser Arg Arg Val Arg Met Gly
1285 1290 1295

Val Phe Gly Cys Trp Leu Ala Phe Ala Val Gly Leu Phe Lys Pro Val
1300 1305 1310

Ser Asp Pro Val Gly Ala Ala Cys Glu Phe Asp Ser Pro Glu Cys Arg
1315 1320 1325

Asn Ile Leu His Ser Phe Glu Leu Leu Lys Pro Trp Asp Pro Val Arg
1330 1335 1340

Ser Leu Val Val Gly Pro Val Gly Leu Gly Leu Ala Ile Leu Gly Arg
1345 1350 1355 1360

Leu Leu Gly Gly Ala Arg Cys Ile Trp His Phe Leu Leu Arg Leu Gly
1365 1370 1375

Ile Val Ala Asp Cys Ile Leu Ala Gly Ala Tyr Val Leu Ser Gln Gly
1380 1385 1390

Arg Cys Lys Lys Cys Trp Gly Ser Cys Ile Arg Thr Ala Pro Asn Glu
1395 1400 1405

Val Ala Phe Asn Val Phe Pro Phe Thr Arg Ala Thr Arg Ser Ser Leu
1410 1415 1420

Ile Asp Leu Cys Asp Arg Phe Cys Ala Pro Lys Gly Met Asp Pro Ile
1425 1430 1435 1440

Phe Leu Ala Thr Gly Trp Arg Gly Cys Trp Ala Gly Arg Ser Pro Ile
1445 1450 1455

Glu Gln Pro Ser Glu Lys Pro Ile Ala Phe Ala Gln Leu Asp Glu Lys
1460 1465 1470

Lys Ile Thr Ala Arg Thr Val Val Ala Gln Pro Tyr Asp Pro Asn Gln
1475 1480 1485

Ala Val Lys Cys Leu Arg Val Leu Gln Ser Gly Gly Arg Trp Trp Leu
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Ser Gly Pro Lys Ser Gly Gln Gly Phe Arg Cys Ser Ile Pro Ser Pro
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Phe Phe Pro Thr Gly Val Lys Val Asp Pro Asp Cys Arg Val Val Val
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Asp Pro Asp Thr Phe Thr Ala Ala Leu Arg Ser Gly Tyr Ser Thr Thr
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Asn Leu Val Leu Gly Val Gly Asp Phe Ala Gln Leu Asn Gly Leu Lys
1555 1560 1565

Ile Arg Gln Ile Ser Lys Pro Ser Gly Gly Pro His Leu Met Ala
1570 1575 1580

Ala Leu His Val Ala Cys Ser Met Ala Leu His Met Leu Ala Gly Ile
1585 1590 1595 1600

Tyr Val Thr Ala Val Gly Ser Cys Gly Thr Gly Thr Asn Asp Pro Trp
1605 1610 1615

Cys Ala Asn Pro Phe Ala Val Pro Gly Tyr Gly Pro Gly Ser Leu Cys
1620 1625 1630

Thr Ser Arg Leu Cys Ile Ser Gln His Gly Leu Thr Leu Pro Leu Thr
1635 1640 1645

Ala Leu Val Ala Gly Phe Gly Ile Gln Glu Ile Ala Leu Val Val Leu
1650 1655 1660

Ile Phe Val Ser Ile Gly Gly Met Ala His Arg Leu Ser Cys Lys Ala
1665 1670 1675 1680

Asp Met Leu Cys Val Leu Leu Ala Ile Ala Ser Tyr Val Trp Val Pro
1685 1690 1695

Leu Thr Trp Leu Leu Cys Val Phe Pro Cys Trp Leu Arg Cys Phe Ser
1700 1705 1710

Leu His Pro Leu Thr Ile Leu Trp Leu Val Phe Phe Leu Ile Ser Val
1715 1720 1725

Asn Met Pro Ser Gly Ile Leu Ala Met Val Leu Leu Val Ser Leu Trp
1730 1735 1740

Leu Leu Gly Arg Tyr Thr Asn Val Ala Gly Leu Val Thr Pro Tyr Asp
1745 1750 1755 1760

Ile His His Tyr Thr Ser Gly Pro Arg Gly Val Ala Ala Leu Ala Thr
1765 1770 1775

Ala Pro Asp Gly Thr Tyr Leu Ala Ala Val Arg Arg Ala Ala Leu Thr
1780 1785 1790

Gly Arg Thr Met Leu Phe Thr Pro Ser Gln Leu Gly Ser Leu Leu Glu
1795 1800 1805

Gly Ala Phe Arg Thr Arg Lys Pro Ser Leu Asn Thr Val Asn Val Ile
1810 1815 1820

Gly Ser Ser Met Gly Ser Gly Gly Val Phe Thr Ile Asp Gly Lys Val
1825 1830 1835 1840

Lys Cys Val Thr Ala Ala His Val Leu Thr Gly Asn Ser Ala Arg Val
1845 1850 1855

Ser Gly Val Gly Phe Asn Gln Met Leu Asp Phe Asp Val Lys Gly Asp
1860 1865 1870

Phe Ala Ile Ala Asp Cys Pro Asn Trp Gln Gly Ala Ala Pro Lys Thr
1875 1880 1885

Gln Phe Cys Thr Asp Gly Trp Thr Gly Arg Ala Tyr Trp Leu Thr Ser
1890 1895 1900

Ser Gly Val Glu Pro Gly Val Ile Gly Lys Gly Phe Ala Phe Cys Phe
1905 1910 1915 1920

Thr Ala Cys Gly Asp Ser Gly Ser Pro Val Ile Thr Glu Ala Gly Glu
1925 1930 1935

Leu Val Gly Val His Thr Gly Ser Asn Lys Gln Gly Gly Ile Val
1940 1945 1950

Thr Arg Pro Ser Gly Gln Phe Cys Asn Val Ala Pro Ile Lys Leu Ser
1955 1960 1965

Glu Leu Ser Glu Phe Phe Ala Gly Pro Lys Val Pro Leu Gly Asp Val
1970 1975 1980

Lys Val Gly Ser His Ile Ile Lys Asp Ile Ser Glu Val Pro Ser Asp
1985 1990 1995 2000

Leu Cys Ala Leu Leu Ala Ala Lys Pro Glu Leu Glu Gly Gly Leu Ser
2005 2010 2015

Thr Val Gln Leu Leu Cys Val Phe Phe Leu Leu Trp Arg Met Met Gly
2020 2025 2030

His Ala Trp Thr Pro Leu Val Ala Val Ser Phe Phe Ile Leu Asn Glu
2035 2040 2045

Val Leu Pro Ala Val Leu Val Arg Ser Val Phe Ser Phe Gly Met Phe
2050 2055 2060

Val Leu Ser Trp Leu Thr Pro Trp Ser Ala Gln Val Leu Met Ile Arg
2065 2070 2075 2080

Leu Leu Thr Ala Ala Leu Asn Arg Asn Arg Trp Ser Leu Ala Phe Phe
2085 2090 2095

Ser Leu Gly Ala Val Thr Gly Phe Val Ala Asp Leu Ala Ala Thr Gln
2100 2105 2110

Gly His Pro Leu Gln Ala Val Met Asn Leu Ser Thr Tyr Ala Phe Leu
2115 2120 2125

Pro Arg Met Met Val Val Thr Ser Pro Val Pro Val Ile Thr Cys Gly
2130 2135 2140

Val Val His Leu Leu Ala Ile Ile Leu Tyr Leu Phe Lys Tyr Arg Gly
2145 2150 2155 2160

Pro His His Ile Leu Val Gly Asp Gly Val Phe Ser Ala Ala Phe Phe
2165 2170 2175

Leu Arg Tyr Phe Ala Glu Gly Lys Leu Arg Glu Gly Val Ser Gln Ser
2180 2185 2190

Cys Gly Met Asn His Glu Ser Leu Thr Gly Ala Leu Ala Met Arg Leu
2195 2200 2205

Asn Asp Glu Asp Leu Asp Phe Leu Met Lys Trp Thr Asp Phe Lys Cys
2210 2215 2220

Phe Val Ser Ala Ser Asn Met Arg Asn Ala Ala Gly Gln Phe Ile Glu
2225 2230 2235 2240

Ala Ala Tyr Ala Lys Ala Leu Arg Val Glu Leu Ala Gln Leu Val Gln
2245 2250 2255

Val Asp Lys Val Arg Gly Thr Leu Ala Lys Leu Glu Ala Phe Ala Asp
2260 2265 2270

Thr Val Ala Pro Gln Leu Ser Pro Gly Asp Ile Val Val Ala Leu Gly
2275 2280 2285

His Thr Pro Val Gly Ser Ile Phe Asp Leu Lys Val Gly Ser Thr Lys
2290 2295 2300

His Thr Leu Gln Ala Ile Glu Thr Arg Val Leu Ala Gly Ser Lys Met
2305 2310 2315 2320

Thr Val Ala Arg Val Val Asp Pro Thr Pro Thr Pro Pro Ala Pro
2325 2330 2335

Val Pro Ile Pro Leu Pro Pro Lys Val Leu Glu Asn Gly Pro Asn Ala
2340 2345 2350

Trp Gly Asp Glu Asp Arg Leu Asn Lys Lys Lys Arg Arg Arg Met Glu
2355 2360 2365

Ala Leu Gly Ile Tyr Val Met Gly Gly Lys Lys Tyr Gln Lys Phe Trp
2370 2375 2380

Asp Lys Asn Ser Gly Asp Val Phe Tyr Glu Glu Val His Asn Asn Thr
2385 2390 2395 2400

Asp Glu Trp Glu Cys Leu Arg Val Gly Asp Pro Ala Asp Phe Asp Pro
2405 2410 2415

Glu Lys Gly Thr Leu Cys Gly His Val Thr Ile Glu Asn Lys Ala Tyr
2420 2425 2430

His Val Tyr Thr Ser Pro Ser Gly Lys Lys Phe Leu Val Pro Val Asn
2435 2440 2445

Pro Glu Asn Gly Arg Val Gln Trp Glu Ala Ala Lys Leu Ser Val Glu
2450 2455 2460

Gln Ala Leu Gly Met Met Asn Val Asp Gly Glu Leu Thr Ala Lys Glu
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Glu Gln Cys Leu Asn Cys
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<210> 2
<211> 1457
<212> PRT
<213> Porcine reproductive and respiratory syndrome virus

<400> 2
Leu Ala Ala Ser Asp Leu Thr Arg Cys Gly Arg Gly Leu Val Val

1

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10

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Thr Glu Thr Ala Val Lys Ile Val Lys Phe His Asn Arg Thr Phe Thr
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Leu Gly Pro Val Asn Leu Lys Val Ala Ser Glu Val Glu Leu Lys Asp
35 40 45

Ala Val Glu His Asn Gln His Pro Val Ala Arg Pro Ile Asp Gly Gly
50 55 60

Val Val Leu Leu Arg Ser Ala Val Pro Ser Leu Ile Asp Val Leu Ile
65 70 75 80

Ser Gly Ala Asp Ala Ser Pro Lys Leu Leu Ala His His Gly Pro Gly
85 90 95

Asn Thr Gly Ile Asp Gly Thr Leu Trp Asp Phe Glu Ser Glu Ala Thr
100 105 110

Lys Glu Glu Val Ala Leu Ser Ala Gln Ile Ile Gln Ala Cys Asp Ile
115 120 125

Arg Arg Gly Asp Ala Pro Glu Ile Gly Leu Pro Tyr Lys Leu Tyr Pro
130 135 140

Val Arg Gly Asn Pro Glu Arg Val Lys Gly Val Leu Gln Asn Thr Arg
145 150 155 160

Phe Gly Asp Ile Pro Tyr Lys Thr Pro Ser Asp Thr Gly Ser Pro Val
165 170 175

His Ala Ala Ala Cys Leu Thr Pro Asn Ala Thr Pro Val Thr Asp Gly
180 185 190

Arg Ser Val Leu Ala Thr Thr Met Pro Pro Gly Phe Glu Leu Tyr Val
195 200 205

Pro Thr Ile Pro Ala Ser Val Leu Asp Tyr Leu Asp Ser Arg Pro Asp
210 215 220

Cys Pro Lys Gln Leu Thr Glu His Gly Cys Glu Asp Ala Ala Leu Lys
225 230 235 240

Asp Leu Ser Lys Tyr Asp Leu Ser Thr Gln Gly Phe Val Leu Pro Gly
245 250 255

Val Leu Arg Leu Val Arg Lys Tyr Leu Phe Ala His Val Gly Lys Cys
260 265 270

Pro Pro Val His Arg Pro Ser Thr Tyr Pro Ala Lys Asn Ser Met Ala
275 280 285

Gly Ile Asn Gly Asn Arg Phe Pro Thr Lys Asp Ile Gln Ser Val Pro
290 295 300

Glu Ile Asp Val Leu Cys Ala Gln Ala Val Arg Glu Asn Trp Gln Thr

305 310 315 320
Val Thr Pro Cys Thr Leu Lys Lys Gln Tyr Cys Gly Lys Lys Lys Thr
325 330 335
Arg Thr Ile Leu Gly Thr Asn Asn Phe Ile Ala Leu Ala His Arg Ala
340 345 350
Val Leu Ser Gly Val Thr Gln Gly Phe Met Lys Lys Ala Phe Asn Ser
355 360 365
Pro Ile Ala Leu Gly Lys Asn Lys Phe Lys Glu Leu Gln Thr Pro Val
370 375 380
Leu Gly Arg Cys Leu Glu Ala Asp Leu Ala Ser Cys Asp Arg Ser Thr
385 390 395 400
Pro Ala Ile Val Arg Trp Phe Ala Ala Asn Leu Leu Tyr Glu Leu Ala
405 410 415
Cys Ala Glu Glu His Leu Pro Ser Tyr Val Leu Asn Cys Cys His Asp
420 425 430
Leu Leu Val Thr Gln Ser Gly Ala Val Thr Lys Arg Gly Gly Leu Ser
435 440 445
Ser Gly Asp Pro Ile Thr Ser Val Ser Asn Thr Ile Tyr Ser Leu Val
450 455 460
Ile Tyr Ala Gln His Met Val Leu Ser Tyr Phe Lys Ser Gly His Pro
465 470 475 480
His Gly Leu Leu Phe Leu Gln Asp Gln Leu Lys Phe Glu Asp Met Leu
485 490 495
Lys Val Gln Pro Leu Ile Val Tyr Ser Asp Asp Leu Val Leu Tyr Ala
500 505 510
Glu Ser Pro Thr Met Pro Asn Tyr His Trp Trp Val Glu His Leu Asn
515 520 525
Leu Met Leu Gly Phe Gln Thr Asp Pro Lys Lys Thr Ala Ile Thr Asp
530 535 540
Ser Pro Ser Phe Leu Gly Cys Arg Ile Ile Asn Gly Arg Gln Leu Val
545 550 555 560
Pro Asn Arg Asp Arg Ile Leu Ala Ala Leu Ala Tyr His Met Lys Ala
565 570 575
Ser Asn Val Ser Glu Tyr Tyr Ala Ser Ala Ala Ala Ile Leu Met Asp
580 585 590
Ser Cys Ala Cys Leu Glu Tyr Asp Pro Glu Trp Phe Glu Glu Leu Val
595 600 605
Val Gly Ile Ala Gln Cys Ala Arg Lys Asp Gly Tyr Ser Phe Pro Gly

610 615 620

Thr Pro Phe Phe Met Ser Met Trp Glu Lys Leu Arg Ser Asn Tyr Glu
625 630 635 640

Gly Lys Lys Ser Arg Val Cys Gly Tyr Cys Gly Ala Pro Ala Pro Tyr
645 650 655

Ala Thr Ala Cys Gly Leu Asp Val Cys Ile Tyr His Thr His Phe His
660 665 670

Gln His Cys Pro Val Thr Ile Trp Cys Gly His Pro Ala Gly Ser Gly
675 680 685

Ser Cys Ser Glu Cys Lys Ser Pro Val Gly Lys Gly Thr Ser Pro Leu
690 695 700

Asp Glu Val Leu Glu Gln Val Pro Tyr Lys Pro Pro Arg Thr Val Ile
705 710 715 720

Met His Val Glu Gln Gly Leu Thr Pro Leu Asp Pro Gly Arg Tyr Gln
725 730 735

Thr Arg Arg Gly Leu Val Ser Val Arg Arg Gly Ile Arg Gly Asn Glu
740 745 750

Val Gly Leu Pro Asp Gly Asp Tyr Ala Ser Thr Ala Leu Leu Pro Thr
755 760 765

Cys Lys Glu Ile Asn Met Val Ala Val Ala Ser Asn Val Leu Arg Ser
770 775 780

Arg Phe Ile Ile Gly Pro Pro Gly Ala Gly Lys Thr Tyr Trp Leu Leu
785 790 795 800

Gln Gln Val Gln Asp Gly Asp Val Ile Tyr Thr Pro Thr His Gln Thr
805 810 815

Met Leu Asp Met Ile Arg Ala Leu Gly Thr Cys Arg Phe Asn Val Pro
820 825 830

Ala Gly Thr Thr Leu Gln Phe Pro Val Pro Ser Arg Thr Gly Pro Trp
835 840 845

Val Arg Ile Leu Ala Gly Gly Trp Cys Pro Gly Lys Asn Ser Phe Leu
850 855 860

Asp Glu Ala Ala Tyr Cys Asn His Leu Asp Val Leu Arg Leu Leu Ser
865 870 875 880

Lys Thr Thr Leu Thr Cys Leu Gly Asp Phe Lys Gln Leu His Pro Val
885 890 895

Gly Phe Asp Ser His Cys Tyr Val Phe Asp Ile Met Pro Gln Thr Gln
900 905 910

Leu Lys Thr Ile Trp Arg Phe Gly Gln Asn Ile Cys Asp Ala Ile Gln

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Pro Asp Tyr Arg Asp Lys Leu Met Ser Met Val Asn Thr Thr Arg Val		
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Thr Tyr Val Glu Lys Pro Val Arg Tyr Gly Gln Val Leu Thr Pro Tyr		
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960		
His Arg Asp Arg Glu Asp Asp Ala Ile Thr Ile Asp Ser Ser Gln Gly		
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Ala Thr Phe Asp Val Val Thr Leu His Leu Pro Thr Lys Asp Ser Leu		
980	985	990
Asn Arg Gln Arg Ala Leu Val Ala Ile Thr Arg Ala Arg His Ala Ile		
995	1000	1005
Phe Val Tyr Asp Pro His Arg Gln Leu Gln Gly Leu Phe Asp Leu Pro		
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Ala Lys Gly Thr Pro Val Asn Leu Ala Val His Cys Asp Gly Gln Leu		
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1040		
Ile Val Leu Asp Arg Asn Asn Lys Glu Cys Thr Val Ala Gln Ala Leu		
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Gly Asn Gly Asp Lys Phe Arg Ala Thr Asp Lys Arg Val Val Asp Ser		
1060	1065	1070
Leu Arg Ala Ile Cys Ala Asp Leu Glu Gly Ser Ser Ser Pro Leu Pro		
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Lys Val Ala His Asn Leu Gly Phe Tyr Phe Ser Pro Asp Leu Thr Gln		
1090	1095	1100
Phe Ala Lys Leu Pro Val Glu Leu Ala Pro His Trp Pro Val Val Ser		
1105	1110	1115
1120		
Thr Gln Asn Asn Glu Lys Trp Pro Asp Arg Leu Val Ala Ser Leu Arg		
1125	1130	1135
Pro Ile His Lys Tyr Ser Arg Ala Cys Ile Gly Ala Gly Tyr Met Val		
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Gly Pro Ser Val Phe Leu Gly Thr Pro Gly Val Val Ser Tyr Tyr Leu		
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Thr Lys Phe Val Lys Gly Gly Ala Gln Val Leu Pro Glu Thr Val Phe		
1170	1175	1180
Ser Thr Gly Arg Ile Glu Val Asp Cys Arg Glu Tyr Leu Asp Asp Arg		
1185	1190	1195
1200		
Glu Arg Glu Val Ala Ala Ser Leu Pro His Gly Phe Ile Gly Asp Val		
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Lys Gly Thr Thr Val Gly Gly Cys His His Val Thr Ser Arg Tyr Leu		

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Pro Arg Val Leu Pro Lys Glu Ser Val Ala Val Val Gly Val Ser Ser		
1235	1240	1245
Pro Gly Lys Ala Ala Lys Ala Leu Cys Thr Leu Thr Asp Val Tyr Leu		
1250	1255	1260
Pro Asp Leu Glu Ala Tyr Leu His Pro Glu Thr Gln Ser Lys Cys Trp		
1265	1270	1275
Lys Met Met Leu Asp Phe Lys Glu Val Arg Leu Met Val Trp Lys Asp		
1285	1290	1295
Lys Thr Ala Tyr Phe Gln Leu Glu Gly Arg Tyr Phe Thr Trp Tyr Gln		
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Leu Ala Ser Tyr Ala Ser Tyr Ile Arg Val Pro Val Asn Ser Thr Val		
1315	1320	1325
Tyr Leu Asp Pro Cys Met Gly Pro Ala Leu Cys Asn Arg Arg Val Val		
1330	1335	1340
Gly Ser Thr His Trp Gly Ala Asp Leu Ala Val Thr Pro Tyr Asp Tyr		
1345	1350	1355
Gly Ala Lys Ile Ile Leu Ser Ser Ala Tyr His Gly Glu Met Pro Pro		
1365	1370	1375
Gly Tyr Lys Ile Leu Ala Cys Ala Glu Phe Ser Leu Asp Asp Pro Val		
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Lys Tyr Lys His Thr Trp Gly Phe Glu Ser Asp Thr Ala Tyr Leu Tyr		
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Glu Phe Thr Gly Asn Gly Glu Asp Trp Glu Asp Tyr Asn Asp Ala Phe		
1410	1415	1420
Arg Ala Arg Gln Glu Gly Lys Ile Tyr Lys Ala Thr Ala Thr Ser Leu		
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Lys Phe Tyr Phe Pro Pro Gly Pro Val Ile Glu Pro Thr Leu Gly Leu		
1445	1450	1455

Asn

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<210> 3
<211> 256
<212> PRT
<213> Porcine reproductive and respiratory syndrome virus

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<400> 3
Met Lys Trp Gly Pro Cys Lys Ala Phe Leu Thr Lys Leu Ala Asn Phe
1 5 10 15

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Leu	Trp	Met	Leu	Ser	Arg	Ser	Ser	Trp	Cys	Pro	Leu	Leu	Ile	Ser	Leu
20								25					30		
Tyr	Phe	Trp	Pro	Phe	Cys	Leu	Ala	Ser	Pro	Ser	Pro	Val	Gly	Trp	Trp
35						40						45			
Ser	Phe	Ala	Ser	Asp	Trp	Phe	Ala	Pro	Arg	Tyr	Ser	Val	Arg	Ala	Leu
50				55						60					
Pro	Phe	Thr	Leu	Ser	Asn	Tyr	Arg	Arg	Ser	Tyr	Glu	Ala	Phe	Leu	Ser
65						70				75			80		
Gln	Cys	Gln	Val	Asp	Ile	Pro	Thr	Trp	Gly	Thr	Lys	His	Pro	Leu	Gly
				85					90			95			
Met	Leu	Trp	His	His	Lys	Val	Ser	Thr	Leu	Ile	Asp	Glu	Met	Val	Ser
					100			105				110			
Arg	Arg	Met	Tyr	Arg	Ile	Met	Glu	Lys	Ala	Gly	Gln	Ala	Ala	Trp	Lys
					115		120				125				
Gln	Val	Val	Ser	Glu	Ala	Thr	Leu	Ser	Arg	Ile	Ser	Ser	Leu	Asp	Val
		130				135				140					
Val	Ala	His	Phe	Gln	His	Leu	Ala	Ala	Ile	Glu	Ala	Glu	Thr	Cys	Lys
		145				150				155			160		
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